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KLOPPROGGE, CORINNA
SCHRODER, HARTWIG
HAEFNER, STEFAN

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<130> BGI-186US

<140> 10/582,918

<141> 2006-06-14

<150> PCT/EP04/014266

<151> 2004-12-15

<150> DE 10359594.5

<151> 2003-12-18

<160> 42

<170> PatentIn Ver. 3.3

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<213> Corynebacterium glutamicum

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<212> DNA

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<223> Description of Artificial Sequence: Synthetic oligonucleotide

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<220>
 <223> Description of Artificial Sequence: Synthetic
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 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 primer

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<210> 18
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<220>
 <223> Description of Artificial Sequence: Synthetic
 primer

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<210> 19
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 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 primer

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<210> 20
 <211> 18
 <212> DNA
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<220>
 <223> Description of Artificial Sequence: Synthetic
 primer

<400> 20
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<210> 21
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<220>
 <223> Description of Artificial Sequence: Synthetic
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 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 primer

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<210> 23
 <211> 34
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 primer

<400> 23
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<210> 24
 <211> 4323
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 plasmid

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<220>
 <223> Description of Artificial Sequence: Synthetic
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<211> 38

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic primer

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38

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<211> 38

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic primer

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38

<210> 28

<211> 1263

<212> DNA

<213> Corynebacterium glutamicum

<400> 28

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1263

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<211> 5860

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
plasmid

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 Leu Val Arg Val Thr Glu Ala Ala Ala Leu Ala Ser Gly Arg Trp Val
 20 25 30
 Gly Arg Gly Met Lys Asn Glu Gly Asp Gly Ala Ala Val Asp Ala Met
 35 40 45
 Arg Gln Leu Ile Asn Ser Val Thr Met Lys Gly Val Val Val Ile Gly
 50 55 60
 Glu Gly Glu Lys Asp Glu Ala Pro Met Leu Tyr Asn Gly Glu Glu Val
 65 70 75 80
 Gly Thr Gly Phe Gly Pro Glu Val Asp Ile Ala Val Asp Pro Val Asp
 85 90 95
 Gly Thr Thr Leu Met Ala Glu Gly Arg Pro Asn Ala Ile Ser Ile Leu
 100 105 110
 Ala Ala Ala Glu Arg Gly Thr Met Tyr Asp Pro Ser Ser Val Phe Tyr
 115 120 125
 Met Lys Lys Ile Ala Val Gly Pro Glu Ala Ala Gly Lys Ile Asp Ile
 130 135 140

Glu	Ala	Pro	Val	Ala	His	Asn	Ile	Asn	Ala	Val	Ala	Lys	Ser	Lys	Gly
145					150					155					160
Ile	Asn	Pro	Ser	Asp	Val	Thr	Val	Val	Val	Leu	Asp	Arg	Pro	Arg	His
				165					170					175	
Ile	Glu	Leu	Ile	Ala	Asp	Ile	Arg	Arg	Ala	Gly	Ala	Lys	Val	Arg	Leu
			180					185					190		
Ile	Ser	Asp	Gly	Asp	Val	Ala	Gly	Ala	Val	Ala	Ala	Ala	Gln	Asp	Ser
		195					200					205			
Asn	Ser	Val	Asp	Ile	Met	Met	Gly	Thr	Gly	Gly	Thr	Pro	Glu	Gly	Ile
		210				215					220				
Ile	Thr	Ala	Cys	Ala	Met	Lys	Cys	Met	Gly	Gly	Glu	Ile	Gln	Gly	Ile
225					230					235					240
Leu	Ala	Pro	Met	Asn	Asp	Phe	Glu	Arg	Gln	Lys	Ala	His	Asp	Ala	Gly
				245					250					255	
Leu	Val	Leu	Asp	Gln	Val	Leu	His	Thr	Asn	Asp	Leu	Val	Ser	Ser	Asp
			260					265					270		
Asn	Cys	Tyr	Phe	Val	Ala	Thr	Gly	Val	Thr	Asn	Gly	Asp	Met	Leu	Arg
		275					280					285			
Gly	Val	Ser	Tyr	Arg	Ala	Asn	Gly	Ala	Thr	Thr	Arg	Ser	Leu	Val	Met
	290					295					300				
Arg	Ala	Lys	Ser	Gly	Thr	Ile	Arg	His	Ile	Glu	Ser	Val	His	Gln	Leu
305					310					315					320
Ser	Lys	Leu	Gln	Glu	Tyr	Ser	Val	Val	Asp	Tyr	Thr	Thr	Ala	Thr	
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